

Nixon, Hargrave, Devans & Doyle LLP

Attorneys & Counselors at Law

Clinton Square
Post Office Box 1051
Rochester, New York 14603-1051
(716) 263-1000

Fax: (716) 263-1600

PRIVILEGE AND CONFIDENTIALITY NOTICE

The information in this fax is intended for the named recipients only. It may contain privileged and confidential matter. If you have received this fax in error, please notify us immediately by a collect telephone call to (716) 263-1000 and return the original to the sender by mail. We will reimburse you for postage. Do not disclose the contents to anyone. Thank you.

FAX

To:	Fax #:	Telephone #:
1) Examiner Ricigliano Group Art Unit 1648	(703) 308-4426	(703) 308-9346
2)		
3)		
4)		
5)		

INTERNATIONAL PHONE NUMBERS MUST INCLUDE COUNTRY & CITY CODE. SEE LOCAL WHITE PAGES FOR CODES NEEDED.

From: Michael L. Goldman	Date: August 26, 1998	No. of Pages: 36 (including this page)	Client/Matter: 19603/461
Comments: FOR EXAMINATION PURPOSES ONLY			

IF YOU DO NOT RECEIVE ALL OF THESE PAGES, PLEASE CONTACT THE FAX OPERATOR AS SOON AS POSSIBLE AT: (716) 263-1660 or 263-1000 (ext. 1660). THANK YOU.

PATENT
Docket No.: 19603/461 (CRF D-1595A)

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicants	:	Barany et al.)	Examiner:
)	J.W. Ricigliano
Serial No.	:	08/794,851)	
)	Art Unit:
Filed	:	February 4, 1997)	1648
)	
For	:	DETECTION OF NUCLEIC ACID)	
		SEQUENCE DIFFERENCES USING THE)	
		LIGASE DETECTION REACTION WITH)	
		ADDRESSABLE ARRAYS)	

SUPPLEMENTAL AMENDMENT

Assistant Commissioner for Patents
Washington, D.C. 20231

Dear Sir:

Please amend the above-identified application as follows:

In the Claims:

Please amend claims 12, 16, 18, 19, 20, 22, 26, 30, 34, and 81 as follows:

12. (Amended) A method according to claim 1, wherein multiple allele differences at two or more [nearby or] adjacent nucleotide positions, or at nucleotide positions which require overlapping oligonucleotide probe sets, in a single target nucleotide sequence or multiple allele differences at two or more [nearby or] adjacent nucleotide positions, or at nucleotide positions which require overlapping oligonucleotide probe sets, in multiple target nucleotide sequences are distinguished with oligonucleotide probe sets having oligonucleotide probes with target-specific portions which overlap.

16. (Amended) A method according to claim 14, wherein multiple allele differences at two or more [nearby or] adjacent nucleotide positions, or at nucleotide positions which require overlapping oligonucleotide probe sets, in a single target nucleotide